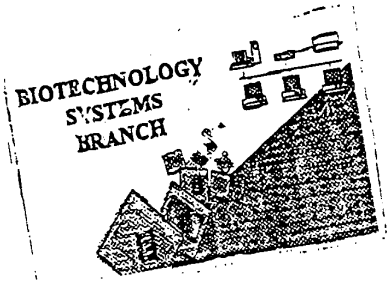

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/601,667
Source: OIPE
Date Processed by STIC: 4/2/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.
FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
   Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED — SUGGESTED CORRECTION — SERIAL NUMBER: 09/601,667

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics / Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to 3; this will prevent "wrapping."

2 ____ Invalid Line Length
The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 ____ Misaligned Amino Numbering
The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 ____ Non-ASCII
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 ✓ Variable Length
Sequence(s) 1 (maybe more) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ Patentin 2.0 "bug"
A "bug" in Patentin version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) _____. Normally, Patentin would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 ____ Skipped Sequences (OLD RULES)
Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 ____ Skipped Sequences (NEW RULES)
Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 ____ Use of n's or Xaa's (NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 ____ Invalid <213> Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 ____ Use of <220>
Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12 ____ Patentin 2.0 "bug"
Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 ____ Misuse of n
n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH – Biotechnology Systems Branch – 08/21/2001

1653

RAW SEQUENCE LISTING DATE: 04/02/2002
PATENT APPLICATION: US/09/601,667 TIME: 15:53:47

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw

Does Not Comply
Corrected Diskette Needed

```
  4 <110> APPLICANT: Morris, Peter
  5       Stiefel, Thomas
  6       Voelter, Wolfgang
  7       Welters, Peter
  9 <120> TITLE OF INVENTION: Recombinant Mistletoe Lectins
 11 <130> FILE REFERENCE: 29841/36636
 13 <140> CURRENT APPLICATION NUMBER: 09/601,667
 14 <141> CURRENT FILING DATE: 2000-10-06
 16 <150> PRIOR APPLICATION NUMBER: PCT/EP99/00696
 17 <151> PRIOR FILING DATE: 1999-02-03
 20 <150> PRIOR APPLICATION NUMBER: D 198 04 210.8
 21 <151> PRIOR FILING DATE: 1998-02-03
 23 <160> NUMBER OF SEQ ID NOS: 39
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 533
 28 <212> TYPE: PRT
C--> 29 <213> ORGANISM: Artificial
 31 <220> FEATURE:
 32 <223> OTHER INFORMATION: mistletoe lectin
 35 <220> FEATURE:
 36 <221> NAME/KEY: SITE
 37 <222> LOCATION: 15
 38 <223> OTHER INFORMATION: product= "Xaa is Asp or Glu"
 39       /label= Xaa1
 41 <220> FEATURE:
 42 <221> NAME/KEY: SITE
 43 <222> LOCATION: 63
 44 <223> OTHER INFORMATION: product= "Xaa is Gly or Gln"
 45       /label= Xaa2
 47 <220> FEATURE:
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 49 <222> LOCATION: 66
 50 <223> OTHER INFORMATION: product= "Xaa is Ile or Val"
 51       /label= Xaa3
 53 <220> FEATURE:
 54 <221> NAME/KEY: SITE
 55 <222> LOCATION: 75
 56 <223> OTHER INFORMATION: product= "Xaa is Leu or Ala"
 57       /label= Xaa4
 59 <220> FEATURE:
 60 <221> NAME/KEY: SITE
 61 <222> LOCATION: 107
 62 <223> OTHER INFORMATION: product= "Xaa is Asp-Arg or
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see pp 1, 5

Xaa can only represent a single amino acid.
Variable length not permitted.
See item 5 on Error Summary Sheet.

RAW SEQUENCE LISTING DATE: 04/02/2002
PATENT APPLICATION: US/09/601,667 TIME: 15:53:47

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw


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63          missing"
64          /label= Xaa5
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69 <223> OTHER INFORMATION: product= "Xaa is Asn or Thr"
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 2 <220> FEATURE:
73 <221> NAME/KEY: SITE
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75 <223> OTHER INFORMATION: product= "Xaa is Pro or Thr"
76          /label= Xaa7
78 <220> FEATURE:
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85 <221> NAME/KEY: SITE
86 <222> LOCATION: 141
87 <223> OTHER INFORMATION: product= "Xaa is Ser or Thr"
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90 <220> FEATURE:
91 <221> NAME/KEY: SITE
92 <222> LOCATION: 145
93 <223> OTHER INFORMATION: product= "Xaa is Phe or Tyr"
94          /label= Xaa10
96 <220> FEATURE:
97 <221> NAME/KEY: SITE
98 <222> LOCATION: 152
99 <223> OTHER INFORMATION: product= "Xaa is Thr or Ala"
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102 <220> FEATURE:
103 <221> NAME/KEY: SITE
104 <222> LOCATION: 177
105 <223> OTHER INFORMATION: product= "Xaa is Ala or Tyr"
106         /label= Xaa12
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111 <223> OTHER INFORMATION: product= "Xaa is Tyr or Asp"
112         /label= Xaa13
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115 <221> NAME/KEY: SITE
116 <222> LOCATION: 185
117 <223> OTHER INFORMATION: product= "Xaa is Ala or Glu"
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120 <220> FEATURE:
121 <221> NAME/KEY: SITE
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/601,667

DATE: 04/02/2002
TIME: 15:53:47

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw

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123 <223> OTHER INFORMATION: product= "Xaa is Val or Met"
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126 <220> FEATURE:
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129 <223> OTHER INFORMATION: product= "Xaa is Ile or Phe"
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132 <220> FEATURE:
133 <221> NAME/KEY: SITE
134 <222> LOCATION: 224
135 <223> OTHER INFORMATION: product= "Xaa is Pro or Ser"
136       /label= Xaa17
138 <220> FEATURE:
139 <221> NAME/KEY: SITE
140 <222> LOCATION: 225
141 <223> OTHER INFORMATION: product= "Xaa is Pro or Thr"
142       /label= Xaa18
144 <220> FEATURE:
145 <221> NAME/KEY: SITE
146 <222> LOCATION: 232
147 <223> OTHER INFORMATION: product= "Xaa is Thr or Ser"
148       /label= Xaa19
150 <220> FEATURE:
151 <221> NAME/KEY: SITE
152 <222> LOCATION: 236
153 <223> OTHER INFORMATION: product= "Xaa is Asp or Ser"
154       /label= Xaa20
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157 <221> NAME/KEY: SITE
158 <222> LOCATION: 287
159 <223> OTHER INFORMATION: product= "Xaa is Asn or Ser"
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164 <222> LOCATION: 290
165 <223> OTHER INFORMATION: product= "Xaa is Cys or Arg"
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169 <221> NAME/KEY: SITE
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171 <223> OTHER INFORMATION: product= "Xaa is Gly or Asn"
172       /label= Xaa23
174 <220> FEATURE:
175 <221> NAME/KEY: SITE
176 <222> LOCATION: 364
177 <223> OTHER INFORMATION: product= "Xaa is Gly or Asp"
178       /label= Xaa24
180 <220> FEATURE:
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RAW SEQUENCE LISTING DATE: 04/02/2002
PATENT APPLICATION: US/09/601,667 TIME: 15:53:47

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw

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162 <222> LOCATION: 426
163 <223> OTHER INFORMATION: product= "Xaa is Gly or Gln"
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166 <220> FEATURE:
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168 <222> LOCATION: 435
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170       /label= Xaa26
172 <220> FEATURE:
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174 <222> LOCATION: 439
175 <223> OTHER INFORMATION: product= "Xaa is Gln or Lys"
176       /label= Xaa27
178 <220> FEATURE:
179 <221> NAME/KEY: SITE
180 <222> LOCATION: 442
181 <223> OTHER INFORMATION: product= "Xaa is Gly or missing"
182       /label= Xaa28
184 <220> FEATURE:
185 <221> NAME/KEY: SITE
186 <222> LOCATION: 443
187 <223> OTHER INFORMATION: product= "Xaa is Arg or Lys"
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190 <220> FEATURE:
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192 <222> LOCATION: 464
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199 <223> OTHER INFORMATION: product= "Xaa is Ala or Gly"
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203 <221> NAME/KEY: SITE
204 <222> LOCATION: 481
205 <223> OTHER INFORMATION: product= "Xaa is Gly or Ala"
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209 <221> NAME/KEY: SITE
210 <222> LOCATION: 483
211 <223> OTHER INFORMATION: product= "Xaa is Ser or Gly"
212       /label= Xaa33
214 <220> FEATURE:
215 <221> NAME/KEY: SITE
216 <222> LOCATION: 484
217 <223> OTHER INFORMATION: product= "Xaa is Gly or Ser"
218       /label= Xaa34
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RAW SEQUENCE LISTING DATE: 04/02/2002
PATENT APPLICATION: US/09/601,667 TIME: 15:53:47

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw

240 <220> FEATURE:
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242 <222> LOCATION: 493
243 <223> OTHER INFORMATION: product= "Xaa is Gly or Tyr"
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249 <223> OTHER INFORMATION: product= "Xaa is Asn or Ser Thr or Lys"
250 /label= Xaa36
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253 <221> NAME/KEY: SITE
254 <222> LOCATION: 501
255 <223> OTHER INFORMATION: product= "Xaa is Ser or Gly"
256 /label= Xaa37
258 <220> FEATURE:
259 <221> NAME/KEY: SITE
260 <222> LOCATION: 502
261 <223> OTHER INFORMATION: product= "Xaa is Leu or Pro"
262 /label= Xaa38
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265 <221> NAME/KEY: SITE
266 <222> LOCATION: 503
267 <223> OTHER INFORMATION: product= "Xaa is Ala or Met"
268 /label= Xaa39
270 <220> FEATURE:
271 <221> NAME/KEY: SITE
272 <222> LOCATION: 504
273 <223> OTHER INFORMATION: product= "Xaa is Met or Val"
274 /label= Xaa40
276 <220> FEATURE:
277 <221> NAME/KEY: SITE
278 <222> LOCATION: 533
279 <223> OTHER INFORMATION: product= "Xaa is Pro or Phe"
280 /label= Xaa41
284 <400> SEQUENCE: 1

W--> 286 Tyr Glu Arg Leu Arg Leu Arg Val Thr His Gln Thr Thr Gly Xaa Glu
287 1 5 10 15
289 Tyr Phe Arg Phe Ile Thr Leu Leu Arg Asp Tyr Val Ser Ser Gly Ser
290 20 25 30
292 Phe Ser Asn Glu Ile Pro Leu Leu Arg Gln Ser Thr Ile Pro Val Ser
293 35 40 45
W--> 295 Asp Ala Gln Arg Phe Val Leu Val Glu Leu Thr Asn Gln Gly Xaa Asp
296 50 55 60
W--> 298 Ser Xaa Thr Ala Ala Ile Asp Val Thr Asn Xaa Tyr Val Val Ala Tyr
299 65 70 75 80
301 Gln Ala Gly Asp Gln Ser Tyr Phe Leu Arg Asp Ala Pro Arg Gly Ala
302 85 90 95
W--> 304 Glu Thr His Leu Phe Thr Gly Thr Thr Arg Xaa Ser Ser Leu Pro Phe

FYI

Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/601,667

DATE: 04/02/2002
TIME: 15:53:48

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw

L:29 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:298 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:3[illegible] M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:3[illegible] M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:328 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:340 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:368 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:393 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:531 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
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L:546 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:549 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:555 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:561 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:564 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:574 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:714 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:726 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:738 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
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L:766 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:879 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:939 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:1002 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:1065 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/601,667

DATE: 04/02/2002
TIME: 15:53:48

Input Set : A:\36636seq.txt
Output Set: N:\CRF3\04022002\I601667.raw

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L:1250 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:1312 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:1342 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:1376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:1[illegible] M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:1[illegible] M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:1453 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:1465 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
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L:1544 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:1581 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
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L:1835 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23
L:1874 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:1913 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:1952 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
L:1991 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
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L:2069 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:2108 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30
L:2147 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
L:2186 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32
L:2226 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33
L:2229 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:33
L:2235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:2242 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:34
L:2245 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:34
L:2251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:2257 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:35
L:2270 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:36
L:2287 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:37
L:2352 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:38
L:2418 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:39